FIGURE 2

1 2	A q	gctgtgcagagccgtctggcagggttgacctcctaaagggatattccatctttattaatcattag 6 	65
66	A 1	tagtgtggtcagagacttagcaccattggtcfcccccaacctggtccagacatttcagcagttta 1	130
	A B	tcggaacagcaacaacagcaacaaaccttcaaaatttacaagtctttaagaaatagaaATGgca tggcttcgcgtgggtggaggaggcgcggctgcaggtctttaagaaatagaaATGgca	195
1		M A	2
196 3		tccaaaagagctctggtcatcctagccaaaggagcagagagatggagacagtgattcctgtgga : S K R A L V I L A K G A E E M E T V I P V D :	260 24
261 362	7	catcatgcggcgagctgggattaaagtcaccgttgcaggcttggctgggaaggaccccgtgcagt I M R R A G I K <u>V T V A G L A G K D P V Q</u> Peptide 1	4.5
326 5 94	حا	gtagccgtgatgtagtgatttgtccggataccagtctggaagaagcaaaacacagggaccatac <u>CSR</u> DVVICPDTSLEEAKTQGPY.	6/
391 201	8	gatgtggttgttcttccaggaggaaatctgggtgcacagaacttatctgagtcggctttggtgaa DVVVLPGGNLGAQNLSESALVK	455 .89
456 103		ggagatcctcaaggagcaggagaacaggaagggcctcatagctgccatctgtgcgggtcctacgg <u>E I L K</u> E Q E N R K G L I A A I C A G P T Peptide 2 *	520 110
521 124	111	ccctgctggctcacgaagtaggctttggatgcaaggttacatcgcacccattggctaaggacaaa A L L A H E V G F G C K V <u>T S H P L A K</u> D K. Peptide 3	585 132
586. 146		atgatgaacggcagtcactacagctactcagagagccgtgtggagaaggacggcctcatcctcac M M N G S H Y S Y S E S R V E K <u>D G L I L T</u> Peptide 4	650 154
651 1 <i>58</i> 1		cageegtgggeetgggaceagettegagtttgegetggeeattgtggaggeaeteagtggeaagg SRGPGTSFEFALAIVEALSGK.	715 175
716 1 8 9	176	acatggctaaccaagtgaaggccccgcttgttctcaaagacTAGagagcccaagccctggaccct DMANQVKAPLVLKD*	780 189
781		ggacccccaggctgagcaggcattggaagcccactagtgtgtccacagcccagtgaacctggcat	845
846		tggaagcccactagtgtgtccacagcccagtgaacctcaggaactaacgtgtgaagtagcccgct	910
911		gctcaggaatctcgccctggctctgtactattctgagccttgctagtagaataaacagttcccca	975



